

## SEQUENCE LISTING

&lt;110&gt; Nunomura, Kiyotada

&lt;120&gt; POLYNUCLEOTIDE AMPLIFICATION METHOD

&lt;130&gt; GP104-02.UT

&lt;160&gt; 9

&lt;170&gt; FastSEQ for Windows Version 3.0

&lt;210&gt; 1

&lt;211&gt; 55

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Primer T7A(-)4190

&lt;400&gt; 1

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55

&lt;210&gt; 2

&lt;211&gt; 19

&lt;212&gt; DNA

&lt;213&gt; Artificial Sequence

&lt;220&gt;

&lt;223&gt; Primer (+)4108

&lt;400&gt; 2

acagcagtagac aaatggcag

19

&lt;210&gt; 3

&lt;211&gt; 8933

&lt;212&gt; RNA

&lt;213&gt; Human Immunodeficiency Virus

&lt;220&gt;

&lt;221&gt; source

&lt;222&gt; (1)...(8933)

&lt;223&gt; Sequence of transcripts produced from the BH10 plasmid.

&lt;400&gt; 3

|             |             |             |             |            |             |      |
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| agggggaaag  | aaaaaaauua  | aaauaaaaaca | uaauaguau   | gcaaggcagg | agcuagaacg  | 240  |
| auucgc当地    | aauccuggcc  | uguuagaaa   | aucagaaggc  | uguagacaaa | uacugggaca  | 300  |
| gcuacaacca  | uccuucagca  | caggauaca   | agaacuuaga  | ucauuauua  | auacaguagc  | 360  |
| aacccucuu   | ugugugcauc  | aaaggauaga  | gauaaaaagac | accaaggaag | cuuuagacaa  | 420  |
| gauagaggaa  | gagcaaaaaca | aaaguaagaa  | aaaagcacag  | caagcagcag | cugacacagg  | 480  |
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| acaucaggcc  | auaucaccua  | gaacuuuaaa  | ugcaugggu   | aaaguaguag | aagagaaggc  | 600  |
| uuucagccca  | gaaguaauac  | ccauguuuuc  | acauuaucu   | gaaggagcca | ccccacaaaga | 660  |
| uuuuaacacc  | augcuaaaca  | cagugggggg  | acaucaagca  | gccaugcaa  | uguuaaaaga  | 720  |
| gaccaucau   | gaggaaagcug | cagaauuggg  | uagaguacau  | ccagugcaug | cagggccuau  | 780  |
| ugcaccaggc  | cagaugagag  | aaccaagggg  | aagugacaua  | gcaggaacua | cuaguacccu  | 840  |
| ucaggaacaa  | auaggaugga  | ugacaaaaua  | uccaccuauc  | ccaguaggag | aaauuuuauaa | 900  |
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| ggacauaaga  | caaggaccaa  | aagaaccuuu  | uagagacaua  | guagaccgg  | ucuauaaaac  | 1020 |
| ucuaagagcc  | gagcaagcuu  | cacaggaggu  | aaaaaaauugg | augacagaaa | ccuuggaccag | 1080 |
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| gcuuucaug   | ccaaguuugu  | uucauaaca   | aagccuuagg  | caucuccuau  | ggcaggaaga  | 5340 |
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| auuacacagg  | ccuguccaaa  | gguauccuuu  | gagccaaauuc | ccauacauua  | uugugccccg  | 6240 |
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<211> 8933  
<212> RNA  
<213> Artificial Sequence

<220>  
<223> Sequence of the IAC-Asrc pseudo target

<221> mutation  
<222> (4135)...(4155)  
<223> Mutated positions: 4135, 4140-1, 4145, 4150,  
4152-3, 4155

<400> 4

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| gagagcguca  | guauuaagcg  | ggggagaauu  | agaucgaugg   | aaaaaaauuc  | gguuaaggcc  | 180  |
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| aaccucuuau  | ugugugcauc  | aaaggauaga  | gauaaaagac   | accaaggaag  | cuuuagacaa  | 420  |
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| ccaaaaugcg  | aaccagagu   | guaagacuu   | uuuuuuaagca  | uugggaccag  | cggcuacacu  | 1140 |
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| uaggaaccaa  | agaaagaugg  | uaaaguguuu  | caauuguggc   | aaagaagggc  | acacagccag  | 1320 |
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<220>

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30

<210> 6

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<220>

<223> Sequence of (-)3837 A30 capture probe.

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67

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<220>

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57

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<220>  
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22

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<213> Artificial Sequence

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4156-57, 4159

<400> 9

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| gagagcguca guauuaagcg ggggagaauu agaucgaugg   | gaaaaaaauuc | gguuaaggcc  | 180  |
| agggggaaag aaaaaauuaa auuuaaaaaca uauaguauugg | gcaagcaggg  | agcuagaacg  | 240  |
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| uuucagccca gaaguauuac ccauguuuuc agcauuaua    | gaaggagcc   | ccccacaaga  | 660  |
| uuuuaacacc augcuuuaaca cagugggggg acaucaagca  | gccaugcaaa  | uguuaaaaaga | 720  |
| gaccaucaau gaggaaagcug cagaauggga uagaguacau  | ccagugcaug  | cagggccuau  | 780  |
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|-------------|-------------|-------------|--------------|-------------|-------------|------|
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| uauiuggcaag | ccaccuggau  | uccugagugg  | gaguuuuguua  | auaccccucc  | uuuagugaaa  | 3180 |
| uuuugguacc  | aguuagagaa  | agaacccaua  | guaggagcag   | aaaccuucua  | uguagauugg  | 3240 |
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| gaagcagaag  | uuauuccagc  | agaaacaggg  | caggaaacag   | cauuuuucu   | uuuuaauua   | 3900 |
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| ccccaaaguc  | aaggaguagu  | agaaucuaug  | aaauaaagaaau | uaaagaaaaaa | uuuaggacag  | 4080 |
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|             |             |             |             |             |             |      |
|-------------|-------------|-------------|-------------|-------------|-------------|------|
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